

SEQUENCE LISTING

<110> Johal, Gurmukh S
Multani, Dilbag S

<120> ISOLATED NUCLEIC ACID MOLECULES ENCODING THE DW3
P-GLYCOPROTEIN OF SORGHUM AND METHODS OF MODIFYING
GROWTH IN TRANSGENIC PLANTS THEREWITH

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<151> 1999-11-12

<150> 09/711,619
<151> 2000-11-13

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<170> PatentIn Ver. 2.1

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 Met Phe Met Lys Gly Phe Ser Gly Asp Leu Glu Ala Ala His Ala Arg

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35	40	45	
gcg ttc aac gcg gag cgc aag atc acg ggg ctg ttc gag gcc aac ctg Ala Phe Asn Ala Glu Arg Lys Ile Thr Gly Leu Phe Glu Ala Asn Leu			192
50	55	60	
cgc ggc ccg ctc cgg cgc tgc ttc tgg aag ggg cag atc gcc ggc agc Arg Gly Pro Leu Arg Arg Cys Phe Trp Lys Gln Ile Ala Gly Ser			240
65	70	75	80
ggc tac ggc gtg gcg cag ttc ctg ctg tac gcg tcc tac gcg ctg ggg Gly Tyr Gly Val Ala Gln Phe Leu Leu Tyr Ala Ser Tyr Ala Leu Gly			288
85	90	95	
ctg tgg tac gcg gcg tgg ctg gtg aag cac ggc gtg tcc gac ttc tcg Leu Trp Tyr Ala Ala Trp Leu Val Lys His Gly Val Ser Asp Phe Ser			336
100	105	110	
cgc acc atc cgc gtg ttc atg gtg ctg atg gtg tcc gcc aac ggc gcc Arg Thr Ile Arg Val Phe Met Val Leu Met Val Ser Ala Asn Gly Ala			384
115	120	125	
gcc gag acg ctg acg ctg gcg ccg gac ttt gtc aag ggc ggg cgc gcg Ala Glu Thr Leu Thr Leu Ala Pro Asp Phe Val Lys Gly Gly Arg Ala			432
130	135	140	
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145	150	155	160
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165	170	175	
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180	185	190	
gtg ttc cgc gac ctg agc ctc cgg gcg cgc gcc ggg aag acg ctg cgc Val Phe Arg Asp Leu Ser Leu Arg Ala Arg Ala Gly Lys Thr Leu Ala			624
195	200	205	
ctg gtg ggt ccg agc ggg tgc ggc aag acg tcg gtg ctg cgc ctg gtg Leu Val Gly Pro Ser Gly Cys Gly Lys Ser Ser Val Leu Ala Leu Val			672
210	215	220	
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225	230	235	240
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245	250	255	

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260						265									270	
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Ala	Tyr	Gly	Arg	Glu	Gly	Ala	Thr	Glu	Ala	Glu	Val	Val	Glu	Ala	Ala	
275						280									285	
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Gln	Arg	Ile	Ile	Ala	Ala	Arg	Ala	Gly	Lys	Gln	Arg	Pro	Ser	Cys	Cys	
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Trp	Thr	Arg	Arg	Pro	Ala	Arg	Trp	Thr	Pro	Ser	Arg	Ser	Gly	Ala	Cys	
340						345									350	
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355						360									365	
ccg	acc	ggc	tgg	cca	ccg	tgc	gct	ccg	ccg	aca	cca	tcg	ccg	tca	tcg	1152
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<213> Sorghum bicolor

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35 40 45

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Arg Gly Pro Leu Arg Arg Cys Phe Trp Lys Gly Gln Ile Ala Gly Ser
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Gly Tyr Gly Val Ala Gln Phe Leu Leu Tyr Ala Ser Tyr Ala Leu Gly
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Leu Trp Tyr Ala Ala Trp Leu Val Lys His Gly Val Ser Asp Phe Ser
100 105 110

Arg Thr Ile Arg Val Phe Met Val Leu Met Val Ser Ala Asn Gly Ala
115 120 125

Ala Glu Thr Leu Thr Leu Ala Pro Asp Phe Val Lys Gly Gly Arg Ala
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Met Arg Ser Val Phe Glu Thr Ile Asp Arg Lys Thr Glu Val Glu Pro
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Asp Asp Val Asp Ala Ala Pro Val Pro Glu Arg Pro Lys Gly Glu Val
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Glu Leu Lys His Val Asp Phe Ser Tyr Pro Ser Arg Pro Asp Ile Gln
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Val Phe Arg Asp Leu Ser Leu Arg Ala Arg Ala Gly Lys Thr Leu Ala
195 200 205

Leu Val Gly Pro Ser Gly Cys Gly Lys Ser Ser Val Leu Ala Leu Val
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Gln Arg Phe Tyr Glu Pro Thr Ser Gly Arg Val Leu Leu Asp Gly Lys
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Asp Val Arg Lys Tyr Asn Leu Arg Ala Leu Arg Arg Val Val Ala Val
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Ala Pro Gln Glu Pro Phe Leu Phe Ala Ala Ser Ile His Asp Asn Ile
260 265 270

Ala Tyr Gly Arg Glu Gly Ala Thr Glu Ala Glu Val Val Glu Ala Ala
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Thr Gln Ala Asn Ala His Arg Phe Ile Ala Ala Leu Pro Glu Gly Tyr
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Gly Thr Gln Val Gly Glu Arg Gly Val Gln Leu Ser Gly Gly Gln Arg
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Gln Arg Ile Ala Ile Ala Arg Ala Gly Lys Gln Arg Pro Ser Cys Cys
325 330 335

Trp Thr Arg Arg Pro Ala Arg Trp Thr Pro Ser Arg Ser Gly Ala Cys

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: oligonucleotide primer designed from sequence of Zea mays Br2 gene

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<213> Sorghum bicolor

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<221> CDS

<222> (1)..(4206)

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Gly Ala Pro His Ala Asp Asp Ala Gly Asp Glu Trp Ala Arg Pro
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gag ctc gag gcc ttc cac ctc ccc tct ccc gcc cac cag cct cct ggc 144
Glu Leu Glu Ala Phe His Leu Pro Ser Pro Ala His Gln Pro Pro Gly
35 40 45

ttc cac cta gcc gct ggg cac caa ccg gaa gct gca gca gag caa ccc 192
Phe His Leu Ala Ala Gly His Gln Pro Glu Ala Ala Glu Gln Pro
50 55 60

acc acg ctc cct gct gcc cgc acc agc gac aca tcc act gct gct 240
Thr Thr Leu Pro Ala Ala Arg Arg Thr Ser Asp Thr Ser Thr Ala Ala
65 70 75 80

ggt gct gct cct tct cct tcg ccg cct ccg ccg gct cct ttg 288
Gly Ala Ala Pro Pro Ser Pro Ser Pro Pro Pro Pro Ala Pro Leu
85 90 95

gag atg gac cag ccg ccc aat gcc aag ccg gcc tcc tcc tcc gcc 336
Glu Met Asp Gln Pro Pro Asn Ala Lys Pro Ala Ser Ser Ser Ala Ala
100 105 110

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Ala Ala Gly Ala Asn Asp Asn Lys Lys Pro Thr Pro Pro Ala Ala Leu
115 120 125

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130 135 140

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Val Gly Thr Leu Gly Ala Leu Val His Gly Cys Ser Leu Pro Val Phe
145 150 155 160

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Leu Arg Phe Phe Ala Asp Leu Val Asp Ser Phe Gly Ser His Ala Asn
165 170 175

gac ccg gac acc atg gtc cgc ctc gtc aag tac gcc ttc tac ttc 576
Asp Pro Asp Thr Met Val Arg Leu Val Val Lys Tyr Ala Phe Tyr Phe
180 185 190

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Leu Val Val Gly Ala Ala Ile Trp Ala Ser Ser Trp Ala Glu Ile Ser
195 200 205

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210	215	220	
tac ctg gac gcg gcg ctg cggt cag gac gtgt tcc ttc ttc gac acc gac Tyr Leu Asp Ala Ala Leu Arg Gln Asp Val Ser Phe Phe Asp Thr Asp			
225	230	235	720
gtgt cgc acc tcg gac gtc atc tac gcc atc aac gcg gac gcc gtgt gtgt Val Arg Thr Ser Asp Val Ile Tyr Ala Ile Asn Ala Asp Ala Val Val			
245	250	255	768
gggt gca gga cgc cat cag cga gaa gct ggg caa cct cat cca cta cat Gly Ala Gly Arg His Gln Arg Glu Ala Gly Gln Pro His Pro Leu His			
260	265	270	816
ggc cac ctt cgt ggc ggg ctt cgt cgt ggg ctt cac cgc cgc ctg gca Gly His Leu Arg Gly Leu Arg Arg Gly Leu His Arg Arg Leu Ala			
275	280	285	864
gct ggc gct cgt cac gct cgc cgt cgt gcc gct cat cgc cgt cat cgg Ala Gly Ala Arg His Ala Arg Arg Ala Ala His Arg Arg His Arg			
290	295	300	912
ggg gct cag cgc cgc gct cgc caa gct ctc ctc cag gag cca gga Gly Ala Gln Arg Arg Ala Arg Gln Ala Leu Leu Gln Glu Pro Gly			
305	310	315	960
cgcc gct gtc ggg cgc cag cgg cat cgc gga gca ggc gct cgc gca gat Arg Ala Val Gly Arg Gln Arg His Arg Gly Ala Gly Ala Arg Ala Asp			
325	330	335	1008
acg gat cgt gca ggc ctt cgt cgg cga gga gcg cga aat gcg ggc gta Thr Asp Arg Ala Gly Leu Arg Arg Gly Ala Arg Asn Ala Gly Val			
340	345	350	1056
ctc ggc ggc gtt ggc cgt cgc gca gaa gat cgg cta ccg cag cgg ctt Leu Gly Val Gly Arg Arg Ala Glu Asp Arg Leu Pro Gln Arg Leu			
355	360	365	1104
cgcc caa ggg gct cgg cct cgg cgg cac cta ctt cac cgt ctt ctg ctg Arg Gln Gly Ala Arg Pro Arg Arg His Leu Leu His Arg Leu Leu Leu			
370	375	380	1152
cta cgg cct cct gct ctg gta cgg cgg aca cct cgt ccg cgg aac cac Leu Arg Pro Pro Ala Leu Val Arg Arg Thr Pro Arg Pro Arg Asn His			
385	390	395	1200
acc aac gga ggg ctc gcc atc gcc acc atg ttc tcc gtc atg atc ggc Thr Asn Gly Gly Leu Ala Ile Ala Thr Met Phe Ser Val Met Ile Gly			
405	410	415	1248
ggg ctg gcc ctc ggg cag tcg gcg ccg agc atg gcc gcg ttc gcc aag Gly Leu Ala Leu Gly Gln Ser Ala Pro Ser Met Ala Ala Phe Ala Lys			
420	425	430	1296
gccc cgc gtgt gcg gcc aag atc ttc cgc atc atc gac cac agg ccg Ala Arg Val Ala Ala Ala Lys Ile Phe Arg Ile Ile Asp His Arg Pro			
435	440	445	1344

ggc atc tcc tcg cg	gac gac ggc gag gac ggc ggc gtg gag ctg gag	1392
Gly Ile Ser Ser Arg Asp Gly Glu Asp Gly Gly Val Glu Leu Glu		
450	455	460
tcg gtg acg ggg cg	gtg gag atg agg ggc gtg gac ttc gcg tac ccg	1440
Ser Val Thr Gly Arg Val Glu Met Arg Gly Val Asp Phe Ala Tyr Pro		
465	470	475
480		
tcg cgg ccg gac gtc ccc atc ctg cgc ggc ttc tcg ctc agc gtg ccc	1488	
Ser Arg Pro Asp Val Pro Ile Leu Arg Gly Phe Ser Leu Ser Val Pro		
485	490	495
gcc ggc aag acc atc gcg ctg gtg ggc agc tcc ggc tcc ggg aag agc	1536	
Ala Gly Lys Thr Ile Ala Leu Val Gly Ser Ser Gly Ser Gly Lys Ser		
500	505	510
acg gtg gtg tcg ctc ctc gag agg ttc tac gac ccc agc gca ggg caa	1584	
Thr Val Val Ser Leu Leu Glu Arg Phe Tyr Asp Pro Ser Ala Gly Gln		
515	520	525
atc ttg ctg gac ggg cat gat ctc aag tcg ctg aag ctc cgg tgg ctc	1632	
Ile Leu Leu Asp Gly His Asp Leu Lys Ser Leu Lys Leu Arg Trp Leu		
530	535	540
cgg cag cag att ggt ctg gtg agc cag gag ccg acg ctg ttc gcg acg	1680	
Arg Gln Gln Ile Gly Leu Val Ser Gln Glu Pro Thr Leu Phe Ala Thr		
545	550	555
560		
agc atc aag gag aac ctg ctg ggg cgg gac agt cag agt gcg acg	1728	
Ser Ile Lys Glu Asn Leu Leu Gly Arg Asp Ser Gln Ser Ala Thr		
565	570	575
cag gcc gag atg gag gag gcc gcc agg gtg gcc aac gcg cac tcc ttc	1776	
Gln Ala Glu Met Glu Ala Ala Arg Val Ala Asn Ala His Ser Phe		
580	585	590
atc gtc aag ctc ccc gac ggc tac gac acg cag gtt ggg gag cgc ggc	1824	
Ile Val Lys Leu Pro Asp Gly Tyr Asp Thr Gln Val Gly Glu Arg Gly		
595	600	605
ctg cag ctc tcc ggt ggg cag aag cag cgc atc gcc atc gcc cgc gcc	1872	
Leu Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg Ala		
610	615	620
atg ctc aag aac ccc gcc atc ctg ctg gac gag gcc acc agc gcg	1920	
Met Leu Lys Asn Pro Ala Ile Leu Leu Asp Glu Ala Thr Ser Ala		
625	630	635
640		
ctg gac tcc gag tct gag aag ctc gtg cag gag gcg ctg gac cgc ttc	1968	
Leu Asp Ser Glu Ser Glu Lys Leu Val Gln Glu Ala Leu Asp Arg Phe		
645	650	655
atg atc ggg cgc acc acc ctg gtg atc gcg cac agg atg tcc acc atc	2016	
Met Ile Gly Arg Thr Thr Leu Val Ile Ala His Arg Met Ser Thr Ile		
660	665	670

cgc aag gcc gac gtg gtg gcc ctg cag ggc ggc ccc gtc tcc gag		2064
Arg Lys Ala Asp Val Val Ala Val Leu Gln Gly Gly Pro Val Ser Glu		
675	680	685
atg ggc gcg cac gac gag ttg atg gcc aag ggc gag aac ggc act tac		2112
Met Gly Ala His Asp Glu Leu Met Ala Lys Gly Glu Asn Gly Thr Tyr		
690	695	700
gcc aag ttc atc cgc atg cag gag cag gcg cac gag gcg gtc		2160
Ala Lys Phe Ile Arg Met Gln Glu Gln Ala His Glu Ala Ala Phe Val		
705	710	715
aac gcc cgc cgc agc agc gcc agg ccc tcc agc gcc cgc aac tcc gtc		2208
Asn Ala Arg Arg Ser Ser Ala Arg Pro Ser Ser Ala Arg Asn Ser Val		
725	730	735
agc tcg ccc atc atg acg cgc aac tcc tcc tac ggc cgc tcc cca tac		2256
Ser Ser Pro Ile Met Thr Arg Asn Ser Ser Tyr Gly Arg Ser Pro Tyr		
740	745	750
tcc cgc cgc ctc tcc gac ttc tcc acc tcc gac ttc acc ctc tcc atc		2304
Ser Arg Arg Leu Ser Asp Phe Ser Thr Ser Asp Phe Thr Leu Ser Ile		
755	760	765
cac gac ccg cac cac cac cggtt acg atg gcc gac aag cag ctc gcg		2352
His Asp Pro His His His Arg Thr Met Ala Asp Lys Gln Leu Ala		
770	775	780
ttc cgc gcc ggc gcc agc tcc ttc ctc cgc ctc gcc agg atg aac tcg		2400
Phe Arg Ala Gly Ala Ser Ser Phe Leu Arg Leu Ala Arg Met Asn Ser		
785	790	795
ccc gag tgg gcc tac gcg ctc gtc ggc tcc ctg ggc tcc atg gtc tgc		2448
Pro Glu Trp Ala Tyr Ala Leu Val Gly Ser Leu Gly Ser Met Val Cys		
805	810	815
ggc tcc ttc agc gcc atc ttc gcc tac atc ctc agc gcc gtg ctc agc		2496
Gly Ser Phe Ser Ala Ile Phe Ala Tyr Ile Leu Ser Ala Val Leu Ser		
820	825	830
gtc tac tac gcg ccg gac cct cgc tac atg aag cgc gag atc gcc aag		2544
Val Tyr Tyr Ala Pro Asp Pro Arg Tyr Met Lys Arg Glu Ile Ala Lys		
835	840	845
tac tgc tac ctg ctc atc ggc atg tcc tcc gcg ggc ctg ctg ttc aac		2592
Tyr Cys Tyr Leu Leu Ile Gly Met Ser Ser Ala Ala Leu Leu Phe Asn		
850	855	860
acg gtg cag cac gtg ttc tgg gac acg gtc ggc gag aac ctc acg aag		2640
Thr Val Gln His Val Phe Trp Asp Thr Val Gly Glu Asn Leu Thr Lys		
865	870	875
cgt gtg cgc gag aag atg ttc gcc gcc gtg ctc cgc aac gag atc gcc		2688
Arg Val Arg Glu Lys Met Phe Ala Ala Val Leu Arg Asn Glu Ile Ala		
885	890	895
tgg ttc gac gcc gac gag aac gcc agc gcg cgc gtc gcc agg ctc		2736

Trp Phe Asp Ala Asp Glu Asn Ala Ser Ala Arg Val Ala Ala Arg Leu			
900	905	910	
gcg ctc gac gcc cag aac gtg cgc tcc gcc atc ggg gac cgt atc tcc	2784		
Ala Leu Asp Ala Gln Asn Val Arg Ser Ala Ile Gly Asp Arg Ile Ser			
915	920	925	
gtc atc gtc cag aac tcg gcg ctc atg ctc gtc gcc tgc acc ggc ggc	2832		
Val Ile Val Gln Asn Ser Ala Leu Met Leu Val Ala Cys Thr Ala Gly			
930	935	940	
ttc gtc ctc cag tgg cgc ctc gcg ctc gtg ctc ctc gcc gtc ttc ccg	2880		
Phe Val Leu Gln Trp Arg Leu Ala Leu Val Leu Leu Ala Val Phe Pro			
945	950	955	960
ctc gtc gtg gcc gcc acc gtg ctg cag aag atg ttc atg aag ggc ttc	2928		
Leu Val Val Ala Ala Thr Val Leu Gln Lys Met Phe Met Lys Gly Phe			
965	970	975	
tcg ggg gac ctg gag gcc gcg cac gcc agg gcc acg cag atc gcg ggc	2976		
Ser Gly Asp Leu Glu Ala Ala His Ala Arg Ala Thr Gln Ile Ala Gly			
980	985	990	
gag gcc gtg gcc aac ctg cgc acc gtg gcc gcg ttc aac gcg gag cgc	3024		
Glu Ala Val Ala Asn Leu Arg Thr Val Ala Ala Phe Asn Ala Glu Arg			
995	1000	1005	
aag atc acg ggg ctg ttc gag gcc aac ctg cgc ggc ccg ctc cg	3072		
Lys Ile Thr Gly Leu Phe Glu Ala Asn Leu Arg Gly Pro Leu Arg Arg			
1010	1015	1020	
tgc ttc tgg aag ggg cag atc gcc ggc agc ggc tac ggc gtg gcg cag	3120		
Cys Phe Trp Lys Gly Gln Ile Ala Gly Ser Gly Tyr Gly Val Ala Gln			
1025	1030	1035	1040
ttc ctg ctg tac gcg tcc tac gcg ctg ggg ctg tgg tac gcg gcg tgg	3168		
Phe Leu Leu Tyr Ala Ser Tyr Ala Leu Gly Leu Trp Tyr Ala Ala Trp			
1045	1050	1055	
ctg gtg aag cac ggc gtg tcc gac ttc tcg cgc acc atc cgc gtg ttc	3216		
Leu Val Lys His Gly Val Ser Asp Phe Ser Arg Thr Ile Arg Val Phe			
1060	1065	1070	
atg gtg ctg atg gtg tcc gcc aac ggc gcc gag acg ctg acg ctg	3264		
Met Val Leu Met Val Ser Ala Asn Gly Ala Ala Glu Thr Leu Thr Leu			
1075	1080	1085	
gcg ccg gac ttt gtc aag ggc ggg cgc gcg atg cgg tcc gtg ttc gag	3312		
Ala Pro Asp Phe Val Lys Gly Gly Arg Ala Met Arg Ser Val Phe Glu			
1090	1095	1100	
acc atc gac cgg aaa acg gag gtg gag ccc gac gac gtg gac gcg gcg	3360		
Thr Ile Asp Arg Lys Thr Glu Val Glu Pro Asp Asp Val Asp Ala Ala			
1105	1110	1115	1120
ccg gtg ccg gag cgg ccc aag ggc gag gtg gag ctg aag cac gtg gac	3408		
Pro Val Pro Glu Arg Pro Lys Gly Glu Val Glu Leu Lys His Val Asp			

1125	1130	1135	
ttc tcg tac ccg tcg cg ^g ccg gac atc cag gtg ttc cgc gac ctg agc Phe Ser Tyr Pro Ser Arg Pro Asp Ile Gln Val Phe Arg Asp Leu Ser			3456
1140	1145	1150	
ctc cg ^g cg ^c gc ^c g ^{gg} a ^a g ^a c ^c t ^t c ^t g ^g c ^g c ^t g ^g t ^t g ^g t ^t c ^c g ^g a ^a c ^c g ^{gg} Leu Arg Ala Arg Ala Gly Lys Thr Leu Ala Leu Val Gly Pro Ser Gly			3504
1155	1160	1165	
tgc ggc a ^a g ^a c ^c t ^t c ^t g ^g t ^t c ^t g ^g c ^c t ^t g ^g a ^a g ^{gg} c ^{cc} Cys Gly Lys Ser Ser Val Leu Ala Leu Val Gln Arg Phe Tyr Glu Pro			3552
1170	1175	1180	
ac ^g tcc g ^{gg} c ^{gc} g ^{tg} c ^{tc} c ^t g ^{ac} g ^{gc} a ^a g ^{ac} g ^{tg} c ^{gc} a ^a g ^{tg} t ^{ac} a ^{ac} Thr Ser Gly Arg Val Leu Leu Asp Gly Lys Asp Val Arg Lys Tyr Asn			3600
1185	1190	1195	1200
ctg cg ^g ctg cg ^c g ^{tg} c ^{tc} g ^{tg} g ^{cg} g ^{tg} g ^{cg} c ^{cg} c ^{ag} g ^{ag} c ^{cg} t ^{tc} Leu Arg Ala Leu Arg Arg Val Val Ala Val Ala Pro Gln Glu Pro Phe			3648
1205	1210	1215	
ctg ttc g ^{cg} g ^{cg} a ^{gc} a ^{tc} c ^{ac} g ^{ac} a ^{ac} a ^{tc} g ^{cg} t ^{ac} g ^{gg} c ^{gc} g ^{ag} g ^{gc} Leu Phe Ala Ala Ser Ile His Asp Asn Ile Ala Tyr Gly Arg Glu Gly			3696
1220	1225	1230	
gc ^g ac ^g gag g ^{cg} g ^{ag} g ^{tg} g ^{tg} g ^{ag} g ^{cg} ac ^g c ^{ag} g ^{cg} a ^{ac} g ^{cg} c ^{ac} Ala Thr Glu Ala Glu Val Val Glu Ala Ala Thr Gln Ala Asn Ala His			3744
1235	1240	1245	
cg ^g ttc atc g ^{cg} g ^{cg} ctg cg ^c g ^{ag} g ^{gc} t ^{ac} g ^{gg} ac ^g c ^{ag} g ^{tg} g ^{gc} g ^{ag} Arg Phe Ile Ala Ala Leu Pro Glu Gly Tyr Gly Thr Gln Val Gly Glu			3792
1250	1255	1260	
cg ^c g ^{gg} g ^{tg} c ^{ag} ctg tcg gg ^c g ^{gg} c ^{ag} c ^{gg} c ^{ag} c ^{gg} atc g ^{cg} atc g ^{cg} Arg Gly Val Gln Leu Ser Gly Gly Gln Arg Gln Arg Ile Ala Ile Ala			3840
1265	1270	1275	1280
cg ^c g ^{cg} ctg g ^{tg} a ^a g ^c g ^{cg} g ^{cc} atc g ^{tg} ctg ctg g ^{ac} g ^{ag} g ^{cg} acc Arg Ala Leu Val Lys Gln Ala Ala Ile Val Leu Leu Asp Glu Ala Thr			3888
1285	1290	1295	
ac ^g g ^{cg} ctg g ^{ac} g ^{cc} g ^{ag} tcg g ^{ag} cg ^g t ^{gc} g ^{tg} c ^{ag} g ^{ag} g ^{cg} ctg g ^{ag} Ser Ala Leu Asp Ala Glu Ser Glu Arg Cys Val Gln Glu Ala Leu Glu			3936
1300	1305	1310	
cg ^c g ^{cg} g ^{gg} t ^{cc} g ^{gg} c ^{gc} a ^{cc} a ^{cc} atc g ^{tg} g ^{tg} g ^{cg} c ^{ac} c ^{gg} ctg g ^{cc} Arg Ala Gly Ser Gly Arg Thr Thr Ile Val Val Ala His Arg Leu Ala			3984
1315	1320	1325	
ac ^g gt ^g c ^{gc} g ^{gc} g ^{cg} c ^{ac} a ^{cc} atc g ^{cg} g ^{tc} atc g ^{ac} g ^{ac} g ^{gc} a ^a g ^{tg} Thr Val Arg Gly Ala His Thr Ile Ala Val Ile Asp Asp Gly Lys Val			4032
1330	1335	1340	
gc ^g gag c ^a g ^{gg} tcg c ^a c ^c ctg c ^t c ^t a ^a g ^c c ^a c ^t c ^c g ^{cc} g ^{ac} g ^{gg} Ala Glu Gln Gly Ser His Ser His Leu Leu Lys His His Pro Asp Gly			4080
1345	1350	1355	1360

tgc tac gcg cggtatgcctgc cag cgg ctg acg ggc ggg tgc cgc 4128
Cys Tyr Ala Arg Met Leu Gln Leu Gln Arg Leu Thr Gly Gly Cys Arg
1365 1370 1375

gcc cggtccgcggccgtcgtccaa cgg ggc cgc cgc gta gga tgg 4176
Ala Arg Ala Ala Ala Val Val Val Gln Arg Gly Arg Arg Val Gly Trp
1380 1385 1390

atg gat gga tca tgg atg agt ttg gtt cct tgataaa 4213
Met Asp Gly Ser Trp Met Ser Leu Val Pro
1395 1400

<210> 9
<211> 1402
<212> PRT
<213> Sorghum bicolor

<400> 9
Met Ser Thr Asn Asp Pro Asp Glu Ile Arg Ala Arg Val Val Val Leu
1 5 10 15

Gly Ala Pro His Ala Asp Asp Asp Ala Gly Asp Glu Trp Ala Arg Pro
20 25 30

Glu Leu Glu Ala Phe His Leu Pro Ser Pro Ala His Gln Pro Pro Gly
35 40 45

Phe His Leu Ala Ala Gly His Gln Pro Glu Ala Ala Ala Glu Gln Pro
50 55 60

Thr Thr Leu Pro Ala Ala Arg Arg Thr Ser Asp Thr Ser Thr Ala Ala
65 70 75 80

Gly Ala Ala Pro Pro Ser Pro Ser Pro Pro Pro Pro Pro Ala Pro Leu
85 90 95

Glu Met Asp Gln Pro Pro Asn Ala Lys Pro Ala Ser Ser Ser Ala Ala
100 105 110

Ala Ala Gly Ala Asn Asp Asn Lys Lys Pro Thr Pro Pro Ala Ala Leu
115 120 125

Arg Asp Leu Phe Arg Phe Ala Asp Gly Leu Asp Cys Ala Leu Met Leu
130 135 140

Val Gly Thr Leu Gly Ala Leu Val His Gly Cys Ser Leu Pro Val Phe
145 150 155 160

Leu Arg Phe Phe Ala Asp Leu Val Asp Ser Phe Gly Ser His Ala Asn
165 170 175

Asp Pro Asp Thr Met Val Arg Leu Val Val Lys Tyr Ala Phe Tyr Phe
180 185 190

Leu Val Val Gly Ala Ala Ile Trp Ala Ser Ser Trp Ala Glu Ile Ser

195 200 205

Cys Trp Met Trp Thr Gly Glu Arg Gln Ser Thr Arg Met Arg Ile Arg
210 215 220

Tyr Leu Asp Ala Ala Leu Arg Gln Asp Val Ser Phe Phe Asp Thr Asp
225 230 235 240

Val Arg Thr Ser Asp Val Ile Tyr Ala Ile Asn Ala Asp Ala Val Val
245 250 255

Gly Ala Gly Arg His Gln Arg Glu Ala Gly Gln Pro His Pro Leu His
260 265 270

Gly His Leu Arg Gly Gly Leu Arg Arg Gly Leu His Arg Arg Leu Ala
275 280 285

Ala Gly Ala Arg His Ala Arg Arg Arg Ala Ala His Arg Arg His Arg
290 295 300

Gly Ala Gln Arg Arg Arg Ala Arg Gln Ala Leu Leu Gln Glu Pro Gly
305 310 315 320

Arg Ala Val Gly Arg Gln Arg His Arg Gly Ala Gly Ala Arg Ala Asp
325 330 335

Thr Asp Arg Ala Gly Leu Arg Arg Gly Ala Arg Asn Ala Gly Val
340 345 350

Leu Gly Gly Val Gly Arg Arg Ala Glu Asp Arg Leu Pro Gln Arg Leu
355 360 365

Arg Gln Gly Ala Arg Pro Arg Arg His Leu Leu His Arg Leu Leu Leu
370 375 380

Leu Arg Pro Pro Ala Leu Val Arg Arg Thr Pro Arg Pro Arg Asn His
385 390 395 400

Thr Asn Gly Gly Leu Ala Ile Ala Thr Met Phe Ser Val Met Ile Gly
405 410 415

Gly Leu Ala Leu Gly Gln Ser Ala Pro Ser Met Ala Ala Phe Ala Lys
420 425 430

Ala Arg Val Ala Ala Ala Lys Ile Phe Arg Ile Ile Asp His Arg Pro
435 440 445

Gly Ile Ser Ser Arg Asp Gly Glu Asp Gly Gly Val Glu Leu Glu
450 455 460

Ser Val Thr Gly Arg Val Glu Met Arg Gly Val Asp Phe Ala Tyr Pro
465 470 475 480

Ser Arg Pro Asp Val Pro Ile Leu Arg Gly Phe Ser Leu Ser Val Pro
485 490 495

Ala Gly Lys Thr Ile Ala Leu Val Gly Ser Ser Gly Ser Gly Lys Ser

500 505 510

Thr Val Val Ser Leu Leu Glu Arg Phe Tyr Asp Pro Ser Ala Gly Gln
515 520 525

Ile Leu Leu Asp Gly His Asp Leu Lys Ser Leu Lys Leu Arg Trp Leu
530 535 540

Arg Gln Gln Ile Gly Leu Val Ser Gln Glu Pro Thr Leu Phe Ala Thr
545 550 555 560

Ser Ile Lys Glu Asn Leu Leu Leu Gly Arg Asp Ser Gln Ser Ala Thr
565 570 575

Gln Ala Glu Met Glu Glu Ala Ala Arg Val Ala Asn Ala His Ser Phe
580 585 590

Ile Val Lys Leu Pro Asp Gly Tyr Asp Thr Gln Val Gly Glu Arg Gly
595 600 605

Leu Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg Ala
610 615 620

Met Leu Lys Asn Pro Ala Ile Leu Leu Leu Asp Glu Ala Thr Ser Ala
625 630 635 640

Leu Asp Ser Glu Ser Glu Lys Leu Val Gln Glu Ala Leu Asp Arg Phe
645 650 655

Met Ile Gly Arg Thr Thr Leu Val Ile Ala His Arg Met Ser Thr Ile
660 665 670

Arg Lys Ala Asp Val Val Ala Val Leu Gln Gly Gly Pro Val Ser Glu
675 680 685

Met Gly Ala His Asp Glu Leu Met Ala Lys Gly Glu Asn Gly Thr Tyr
690 695 700

Ala Lys Phe Ile Arg Met Gln Glu Gln Ala His Glu Ala Ala Phe Val
705 710 715 720

Asn Ala Arg Arg Ser Ser Ala Arg Pro Ser Ser Ala Arg Asn Ser Val
725 730 735

Ser Ser Pro Ile Met Thr Arg Asn Ser Ser Tyr Gly Arg Ser Pro Tyr
740 745 750

Ser Arg Arg Leu Ser Asp Phe Ser Thr Ser Asp Phe Thr Leu Ser Ile
755 760 765

His Asp Pro His His His Arg Thr Met Ala Asp Lys Gln Leu Ala
770 775 780

Phe Arg Ala Gly Ala Ser Ser Phe Leu Arg Leu Ala Arg Met Asn Ser
785 790 795 800

Pro Glu Trp Ala Tyr Ala Leu Val Gly Ser Leu Gly Ser Met Val Cys

805 810 815

Gly Ser Phe Ser Ala Ile Phe Ala Tyr Ile Leu Ser Ala Val Leu Ser
820 825 830

Val Tyr Tyr Ala Pro Asp Pro Arg Tyr Met Lys Arg Glu Ile Ala Lys
835 840 845

Tyr Cys Tyr Leu Leu Ile Gly Met Ser Ser Ala Ala Leu Leu Phe Asn
850 855 860

Thr Val Gln His Val Phe Trp Asp Thr Val Gly Glu Asn Leu Thr Lys
865 870 875 880

Arg Val Arg Glu Lys Met Phe Ala Ala Val Leu Arg Asn Glu Ile Ala
885 890 895

Trp Phe Asp Ala Asp Glu Asn Ala Ser Ala Arg Val Ala Ala Arg Leu
900 905 910

Ala Leu Asp Ala Gln Asn Val Arg Ser Ala Ile Gly Asp Arg Ile Ser
915 920 925

Val Ile Val Gln Asn Ser Ala Leu Met Leu Val Ala Cys Thr Ala Gly
930 935 940

Phe Val Leu Gln Trp Arg Leu Ala Leu Val Leu Leu Ala Val Phe Pro
945 950 955 960

Leu Val Val Ala Ala Thr Val Leu Gln Lys Met Phe Met Lys Gly Phe
965 970 975

Ser Gly Asp Leu Glu Ala Ala His Ala Arg Ala Thr Gln Ile Ala Gly
980 985 990

Glu Ala Val Ala Asn Leu Arg Thr Val Ala Ala Phe Asn Ala Glu Arg
995 1000 1005

Lys Ile Thr Gly Leu Phe Glu Ala Asn Leu Arg Gly Pro Leu Arg Arg
1010 1015 1020

Cys Phe Trp Lys Gly Gln Ile Ala Gly Ser Gly Tyr Gly Val Ala Gln
1025 1030 1035 1040

Phe Leu Leu Tyr Ala Ser Tyr Ala Leu Gly Leu Trp Tyr Ala Ala Trp
1045 1050 1055

Leu Val Lys His Gly Val Ser Asp Phe Ser Arg Thr Ile Arg Val Phe
1060 1065 1070

Met Val Leu Met Val Ser Ala Asn Gly Ala Ala Glu Thr Leu Thr Leu
1075 1080 1085

Ala Pro Asp Phe Val Lys Gly Gly Arg Ala Met Arg Ser Val Phe Glu
1090 1095 1100

Thr Ile Asp Arg Lys Thr Glu Val Glu Pro Asp Asp Val Asp Ala Ala

1105 1110 1115 1120
Pro Val Pro Glu Arg Pro Lys Gly Glu Val Glu Leu Lys His Val Asp
1125 1130 1135

Phe Ser Tyr Pro Ser Arg Pro Asp Ile Gln Val Phe Arg Asp Leu Ser
1140 1145 1150

Leu Arg Ala Arg Ala Gly Lys Thr Leu Ala Leu Val Gly Pro Ser Gly
1155 1160 1165

Cys Gly Lys Ser Ser Val Leu Ala Leu Val Gln Arg Phe Tyr Glu Pro
1170 1175 1180

Thr Ser Gly Arg Val Leu Leu Asp Gly Lys Asp Val Arg Lys Tyr Asn
1185 1190 1195 1200

Leu Arg Ala Leu Arg Arg Val Val Ala Val Ala Pro Gln Glu Pro Phe
1205 1210 1215

Leu Phe Ala Ala Ser Ile His Asp Asn Ile Ala Tyr Gly Arg Glu Gly
1220 1225 1230

Ala Thr Glu Ala Glu Val Val Glu Ala Ala Thr Gln Ala Asn Ala His
1235 1240 1245

Arg Phe Ile Ala Ala Leu Pro Glu Gly Tyr Gly Thr Gln Val Gly Glu
1250 1255 1260

Arg Gly Val Gln Leu Ser Gly Gly Gln Arg Gln Arg Ile Ala Ile Ala
1265 1270 1275 1280

Arg Ala Leu Val Lys Gln Ala Ala Ile Val Leu Leu Asp Glu Ala Thr
1285 1290 1295

Ser Ala Leu Asp Ala Glu Ser Glu Arg Cys Val Gln Glu Ala Leu Glu
1300 1305 1310

Arg Ala Gly Ser Gly Arg Thr Thr Ile Val Val Ala His Arg Leu Ala
1315 1320 1325

Thr Val Arg Gly Ala His Thr Ile Ala Val Ile Asp Asp Gly Lys Val
1330 1335 1340

Ala Glu Gln Gly Ser His Ser His Leu Leu Lys His His Pro Asp Gly
1345 1350 1355 1360

Cys Tyr Ala Arg Met Leu Gln Leu Gln Arg Leu Thr Gly Gly Cys Arg
1365 1370 1375

Ala Arg Ala Ala Ala Val Val Gln Arg Gly Arg Arg Val Gly Trp
1380 1385 1390

Met Asp Gly Ser Trp Met Ser Leu Val Pro
1395 1400